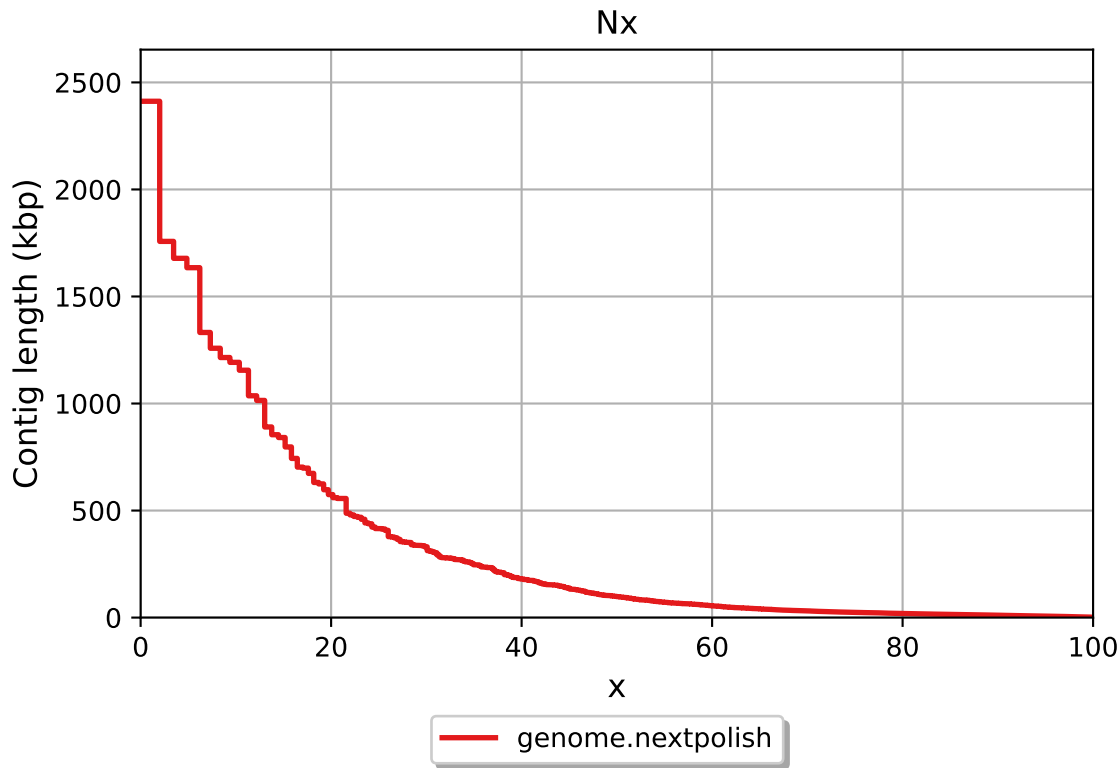


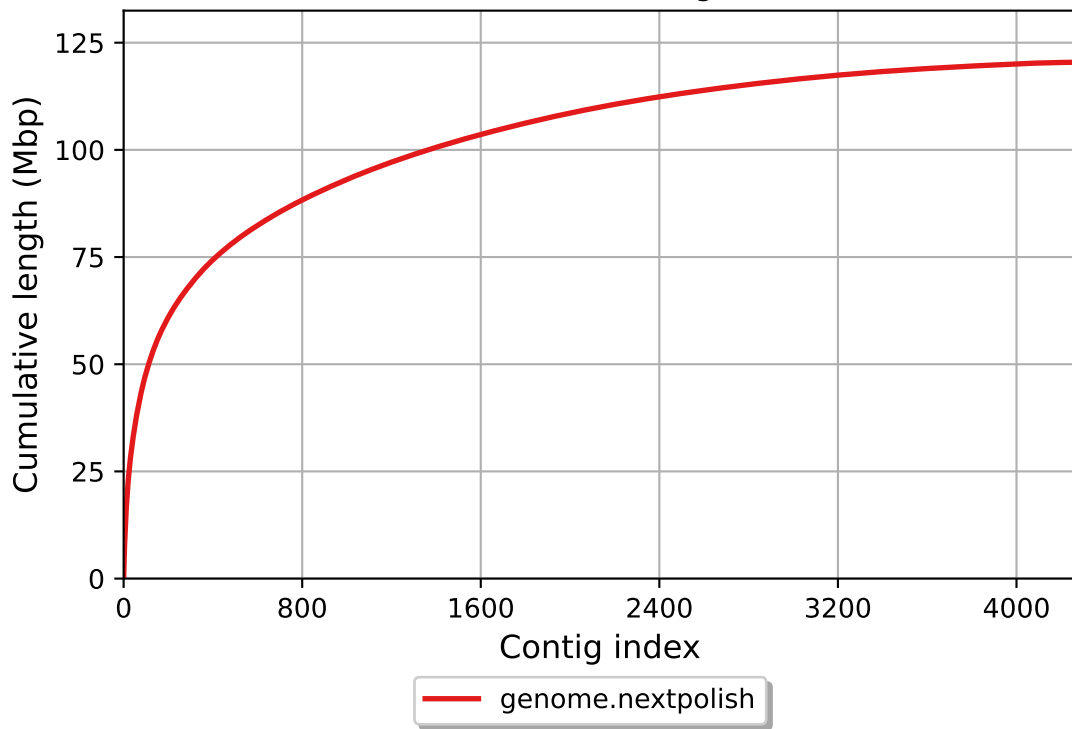
Report

	genome.nextpolish
# contigs (≥ 0 bp)	4277
# contigs (≥ 1000 bp)	4192
# contigs (≥ 5000 bp)	3109
# contigs (≥ 10000 bp)	2122
# contigs (≥ 25000 bp)	845
# contigs (≥ 50000 bp)	390
Total length (≥ 0 bp)	120436059
Total length (≥ 1000 bp)	120383622
Total length (≥ 5000 bp)	116989489
Total length (≥ 10000 bp)	109864775
Total length (≥ 25000 bp)	89461631
Total length (≥ 50000 bp)	73873365
# contigs	4267
Largest contig	2411930
Total length	120433408
GC (%)	45.58
N50	98217
N90	11079
auN	348745.5
L50	193
L90	1983
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	289 + 19 part

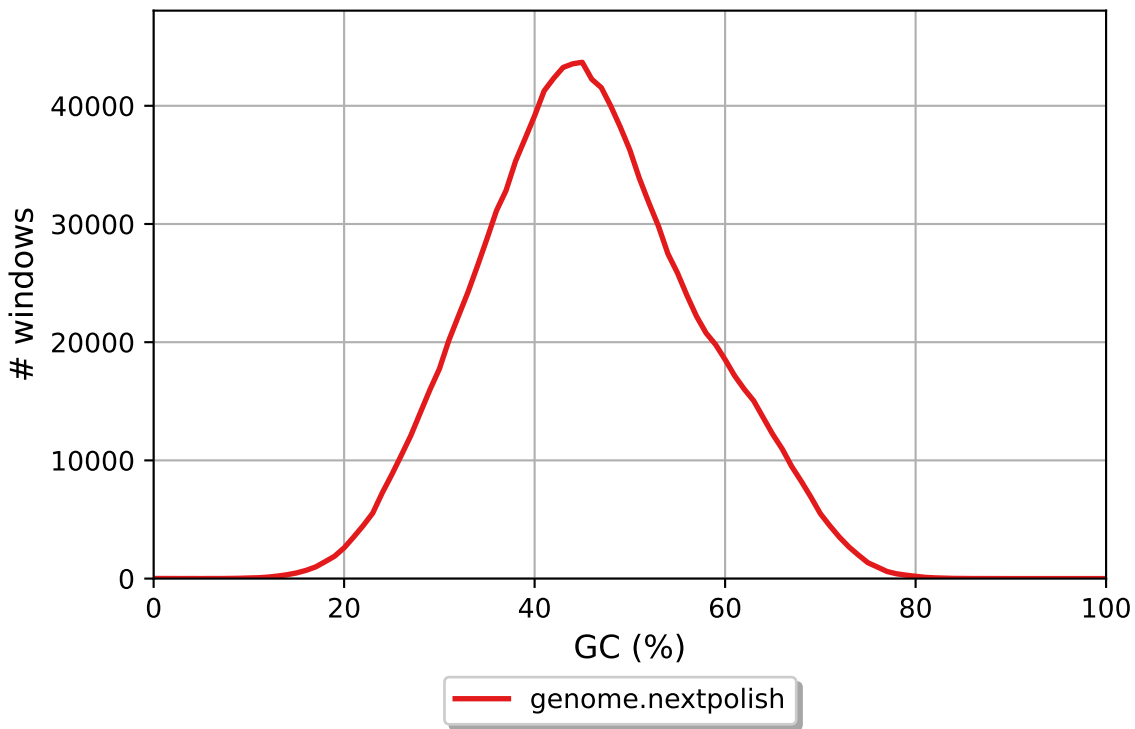
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

